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Fig. 1

EGP-2 promoter sequences

-3967. ^{BglII}AGATCTAGAA TAGAGAGGGA TTGCTGCGAT AGTGGTTAAG GACTTTTACT CTTCAATCTA TATAAAGGAC TTTTGTTC
 -3987. TACTCATCTA TTACTTATGG GATAACAAA ATTTTATAGAA CTGGTAGTCT AATTTATAT ATATATATAT ATATATATAT
 -3987. ATATATATAT ATATATATAT ATATATTTT TTTTATTTT TTTTAGACAG AGTTTGTCTC TTGTTGCCCA GGCTGGAGTG
 -3727. CAATGGCATG ATCTTCGCTC ACCACAACCT CCGCTCCTCG GGTTCAAAGTG ATTCCTCTGC CTCAGCCTCC CAAGTATCTG
 -3647. GAATTACAGG CATGTGCCAC CATGCCACG TAATTTTAT ATTTTATAGTA GAGACAGGTT TTCACACAGGT TGCCACAGGT
 -3567. GCTCTCAAC TCCTGACCTC AAGTGATCCA CCGCTTTGG AATATGGCAC TAGAATACA AAAGATTCAC. AATTAACA TAAACGAGT
 -3487. CTAGCCTGAA AATATTAATA AATGTCTTA AATATGGCAC TAGAATACA AAAGATTCAC. AATTAACA TAAACGAGT
 -3407. AATTTGAGC AAAGATGAC AAATTGAGAA GGTTTAAATG AGTACTAAA ATAAACAATA CCGGCGG(GTGCA)GTGGCTCA p39^B
 -3327. TGCCTGTAAT CCCAGCACTT TGGGAAGCTG ^{SpI}AGGCGGGTGG ATCACCCTGAG GTCAGGAGTT CAAGACCAGC CTGGCCCAACG
 -3247. TAGTGAAC CCGTCTCTAC TAAAAATACA AAAATTAGCC GGCGAGGTG GCAGGCGCT. GTAATCACAG CTAATCGGGA
 -3167. GGCTGAGACA GGAGATTTGC TTGAACCCAG GAGGTGAGG TTGCAGTGAG CTGAGAACAC GCCATTGTAC TCCAGCCTGG
 -3087. GTAACAGAT TGAATCTCTA TCTTAAAAA AAAAAGAGG CGGACACGGT GGCTTGACCC TGTAATCCCA GCACCTTGGG
 -3007. AGCCGAGGC AAGAGGATCA CAAAGTCAGG AGATCAGAC CATCTGGCC AACATGTGTA AACTCTGTCT CAACTGNA
 -2927. TACAAAATTT AGCCGGTGT GTTGGTGGG GCCTGTAATC CCAGCTATTC AGGAGGTGA GGCAGAGAA TTGCTTGAAC
 -2847. CCAAGAGGTG GAGTTGCAG TCCGCCAAGA TCATGCCACT GCATGCAGC TTGGTGACA GAGCAAGACC CCATCTCAA
 -2767. AAAAAAAGG AAGAAAAAAT ACCCTGGATC AGCCGGGTGT GTGGCTCAA GCCTGTAATC CCIAGCACTTTGGAGGTGA p39^{B4-7}
 -2687. GGTGGGCAGA TCACCTGAGG TCAGGAGTTC AAGACCAGCC TGACCAACAT GGAGAACCC CATCTCTACT AAAAATACAA
 -2607. AAAATTAGCC GGACGTGGTG GCACATGCTT GTAATCCAG CTACTCAGGA GGCTGAGGCA GGAGATTCG CTGAATCCGG
 -2547. GAGGCGGAGG TTGTGGTGAG GTGAGATGAT GCCATTGCAC TCCAGCCTGG GCAACAGAG CAAACTCTG CCTCAAAAA
 -2447. AGAAAGAAA AAAAAAAGA AAAAAAGAA AAAATACCT GGATGTATAC TCAGATACAA ^{AP-1}TGAGTCAGAG ATTAGTCTGG
 -2367. TATTTTGTCA TTTATTTAAT AATTATGCTT ACTCAATCA CTT(TATTGTAA)TAACAATA AATAGCTGC CAGTTATAAG p39^{B17-1}
 -2287. AAGATGAGT TCTCCGATT AGTAAACAG AATTAGACT CAGAAATGGAA CATTTTGCCA ATAAAGCCAC AATAACCACT
 -2207. TAGTTTATTC TTGGGAAAG TATATGTAAT TTGGAGAAAG GCAAACTTCC ^{Sp-1}GAAACATC CAAATTCAG CAGACAACAA
 -2127. AAATCTGGT AACTTCTTCC ^{Sp-1}GATTGTGA GTACTATTC(TTTT)TTTTTG TTGTTTGT TTTTTTTTT GAGACGGAGT p39^{B15-1}
 -2047. TTCGCTCTTG TTGCCAGGC TGGAGTGCA TGGCGAATG TTGTTCACT GCAACCTCTG CCTCCAGGT TCAAGTGATT
 -1967. CTCCTGCCTC AGTCTCCTGA GTAGCTGGGA TTACAGGCC ^{Sp-1}CCGACCCAC GCCTGGCTAA CTTCTGTAT TTTTAGTAGA
 -1887. GACGGGTTT CACCATGTTG GCCAGGCTGG TCTCGAATC CTGACCTTAG GTGATCCCGC ^{Sp-1}CGCTCGGCC TCCCAAGTG
 -1807. CTGAGATTAC AGGCATGAGC CACCGTACCT GGCCTAATA CCTATTTC TATACCACGT GAAATTTAAA TTATACAAAA

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Fig. 1, contd.

-1727. CAATATATAG AGGTACTTAG AACAGCATGA CTATTACAT TAATCAACTT GCCGGCACTT CAACAGAATA CAACATAGAA
-1647. ATGATTGTTT TAATATAAC ATAAGCTTTG ATTGACATA TACTTGAGA AATTAATCAA ACTTAGCTGA ATCTTAAAT
-1567. TCGCTTTTTC CTTTTCTT TTTTCTTAT TTTTGTAGT GAGTCTTGT CTGTTGCCAG ACTGGAGTGC
-1487. AGCGTTTGG TCTGGCTCA CCGCAACCTC CGACTCTCTG GTTCAACGGA TTCTCCTGCC TCAGCCTCCT GAGTAGTGG
-1407. GATTACAGGT GCCTGCCACC ACACCTGGCT ACTTTTGTGA TTTTGTAGTTG AGATGGGTTT CACCATGTTG GCCAGGATGG
-1327. TCTCGAATC CTTTTCTT ATCTGCCAC ATCTGCCAC CTGTGCCCC AGCAAGGTGC TGGATTACA AGCATGAGCC ACCGTGCCCA
-1247. GCCTCCTTTC CTTTTCTT CTCTTACTT TATGATTTCT TTAGTGGATA AAAAGCTTTT AAAAATAGG TTACAATGAT
-1167. ATTACAGCTA ACAAAAAATA ACATTAAAAA ACHATAAATA GTATATATAT GAATATTTTA TAATTATTTT AATATTGTAA
-1087. TAATATAGT TGTGTGAT TGAATTCATC TGCACGGAAA TCGATTACTG TCCTTTCTTT CTATTTCCCT ATATTTTCTT p39¹⁷⁻²
-1017. TCCGAAGCT CATCAACAT TTGGTTCTTT AATAGTAACC AAAACCCGAA ATCATCTCG TTCTCAGTAT TTGGCTCTAT
-937. GCGAAGCT TTTCTTTT CTCTTTTCT TTTTCTTTGA GACGGAGTCT TGCTCCTGTC GCCCAGGCTG GAGTGTATG
-857. GCACGATCT TGCTCAGTC AACCTCAGCC TCCCCAGTAG CTGGGATAC AGCATGCGC CACCACGCCG GGCTAATTT
-777. GATCTTTTA GTAGAGCGG CTTTCTTCA TGTGTGTCAG GCTGTCTCG AACTTCAAC CTCAGGTGAT CCGCGCGCT
-697. CGCGCTCCC IAAGTGCTAG ATTACAGCG TGAGCCACCG CGCTACCTT GGAACACCT TTTCTTACAT CTTCAGTGC p39²⁴⁻¹
-617. TAGAATGCT TAGAATAAC AAAAAGAT TATTAGAGT AATTATAAG AATCACTCAT TTTCTTCCA AGAGGCCAA
-537. GATTTCTTCT TTCTTTTCT TTTCTTTCTA ATTTCAAGG AGTATAATTA AATTGCCAGG TAAAGCTGA
-457. AGCTTTT TTATAGTGT CTGGAAGTT CTCTGCTGT GTTTGATTT CTTTAGCT CCACCTTCTT CTATCCAGTT
-377. CCGCACCTT TCCCCCAGG CCCCATTTCT CAAGGCTTCAGAGCAGGCT CCCTCGGTTA IAAGGAGTCT TCAGCACAGA p39⁵¹¹⁻¹
-297. ATCTTCAAC CTCTCTGGAG GCCACCAAG ATCCCTAAC CCGCCATGA GACGAAGCAC CTGCTGCTG GCGGAGCGG
-217. GCGGCGGGC CCACACCTCT AGT GGAGAGGGCC GCGCCCCAAC TGCAGCGCGG GGGTGGGG AGGGAGCTT ACTCACTCCC
-137. CCAACTCCG GCGGCTGACT CAACAGAG CACACAGCGC CAGAGGTGAG CAGTCCCGG AAGGGGCGA GAGGCGGTC
-57. CGCAGGTCCG CAGGTGTC GCTGCTCTCC CGAGCGGAC CCGCTGCGC CAGGCTCTT AGTCTTCTG CGAGCAGACCTTCGAGCC p39^{512-2,3}
+23. GGTCCGGGA CCCCCTGTC GCTGCTCTCC CGAGCGGAC CCGCTGCGC CAGGCTCTT AGTCTTCTT CCGCGCGCGC GCAGCATGTC
+103. TGTCCCACTC CCGCGGACG CCTTCCCGG AGTCCCGG CCTTCCCG CCTCTTCT CCGCGCGCGC GCAGCATGTC
+183. GCGCGGAG GTCTGCTCTT TGGGCTTCT GTTGTGCGG GCGAGCGCA CTTTGCGCG AGCTTCTGGA GGTGAGGCGC
+263. GGAATGAGC AGATTTGTC AGCTGGGCTG GCGTGGCGG CA

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Fig. 2

